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His Phe Ile Asp Glu Gly Arg Asn Ser Gly Gly His Val Phe Asp Tyr		
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Val Leu Glu Asp Cys Thr Val Thr Ile Ser Gln Lys Met Asn Met Asn		
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 Thr Leu Leu Met Glu Asp Leu Asn Asp Gly Leu Arg Thr Pro Gly Ala
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 Ile Met Leu Gly Gly Gly Asn Pro Ala Gln Ile Pro Glu Met Gln Asp
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 Tyr Phe Gln Thr Leu Leu Thr Asp Met Leu Glu Ser Gly Lys Ala Thr
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gat gca ctg tgt aac tac gac ggt cca cag ggg aaa acg gag cta ctc 240
 Asp Ala Leu Cys Asn Tyr Asp Gly Pro Gln Gly Lys Thr Glu Leu Leu
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 Thr Leu Leu Ala Gly Met Leu Arg Glu Lys Leu Gly Trp Asp Ile Glu
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 Pro Gln Asn Ile Ala Leu Thr Asn Gly Ser Gln Ser Ala Phe Phe Tyr
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T3330" 69/9960

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 Cys Gly Ile Ile Ile Ala Asn Glu Lys Ile Ile Thr Ala Ile Thr Asn
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Tyr	Phe 50	Gln	Thr	Leu	Leu	Thr 55	Asp	Met	Leu	Glu	Ser 60	Gly	Lys	Ala	Thr
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Asn 210	Gln	His	Gly	Ile	Pro	Leu 215	Val	Ile	Asp	Asn	Ala 220	Tyr	Gly	Val	Pro
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